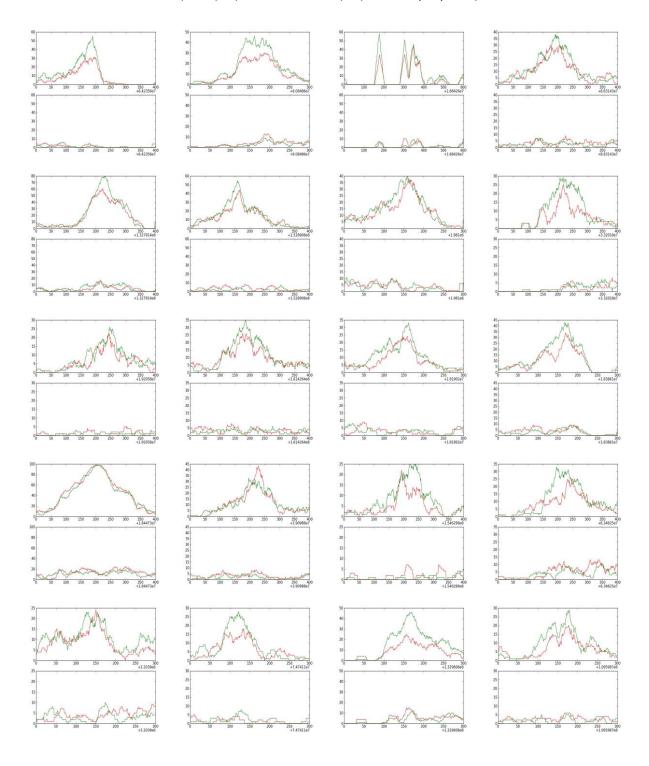
This file contains the peak profile of the top 20 unique peaks in each comparison (NRSF, ATF4 and H3K27me3). The top subplots show the ChIP samples, whereas the bottom subplots show the input samples. In the H3K27me3 data, note that red and orange lines represent the HPV(-) cell lines, blue and cyan lines represent the HPV(+) cell lines (same color coding as in the Figure 4).

Format: (Transcript factor) – (programs compared) –(program-specific unique peaks)

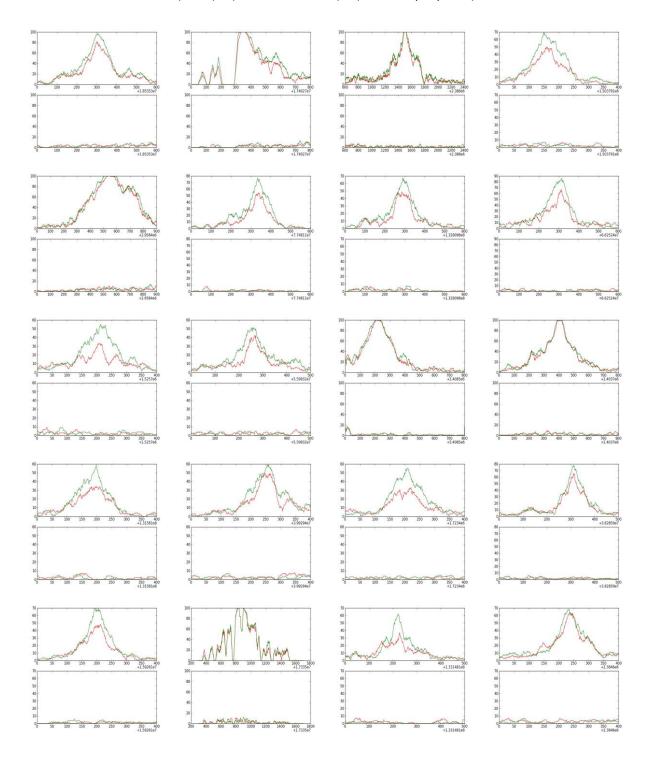
(NRSF) – (PePr vs MACS-CR) – (PePr-unique peaks)



(NRSF) – (PePr vs MACS-CR) – (MACS-CR-unique peaks)



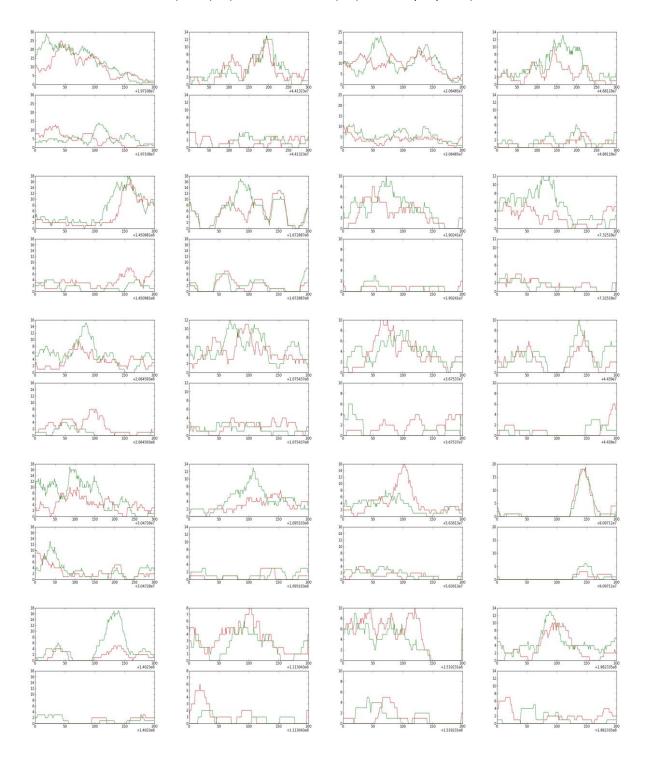
(NRSF) – (PePr vs MACS-SA) – (PePr-unique peaks)



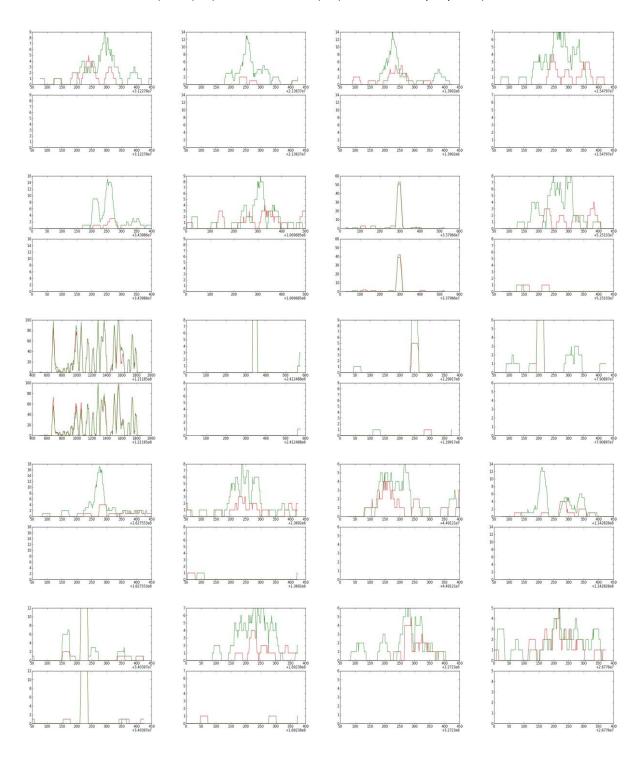
(NRSF) – (PePr vs MACS-SA) – (MACS-SA-unique peaks)



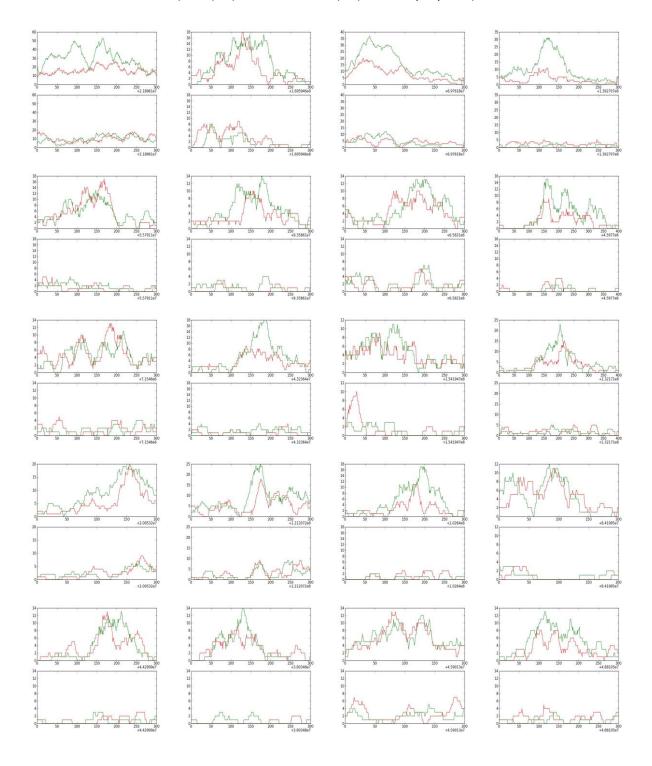
(NRSF) – (PePr vs ZINBA-CR) – (PePr-unique peaks)



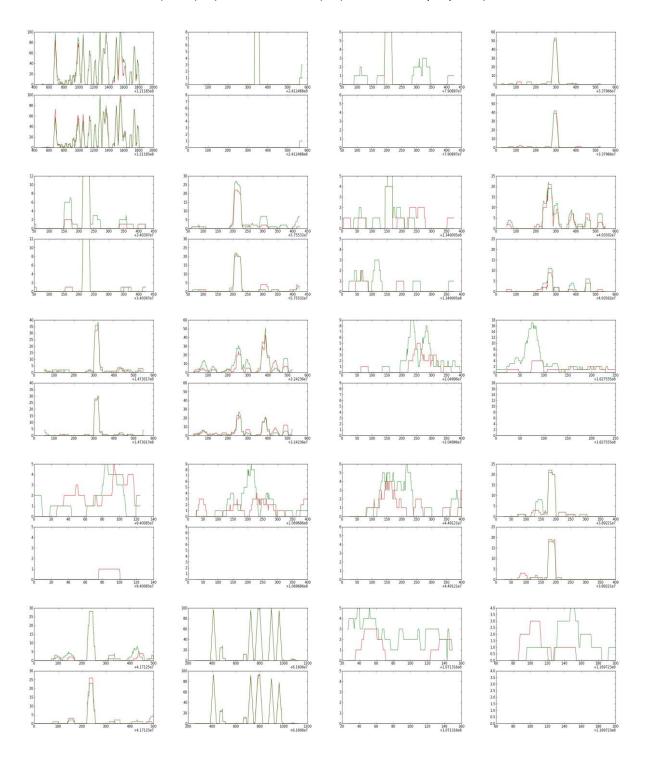
(NRSF) – (PePr vs ZINBA-CR) – (ZINBA-CR-unique peaks)



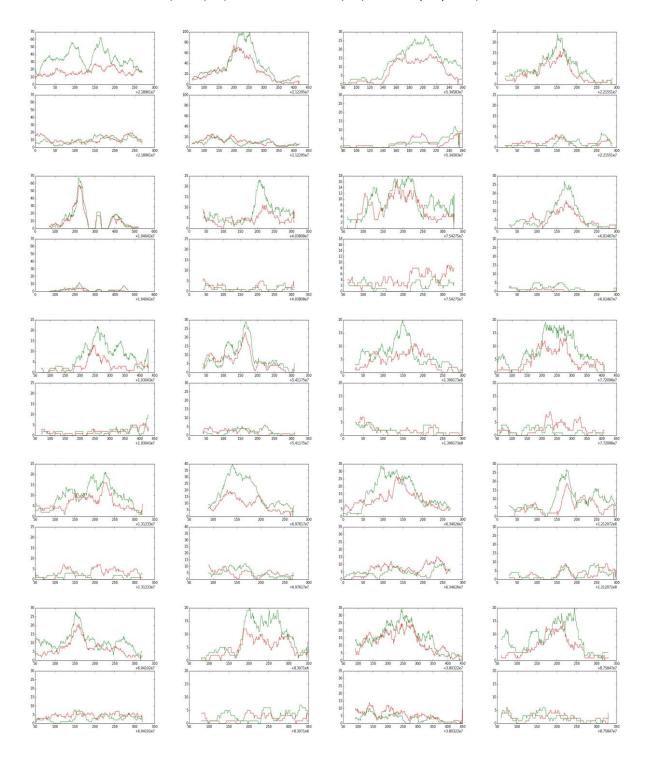
(NRSF) – (PePr vs ZINBA-SA) – (PePr-unique peaks)



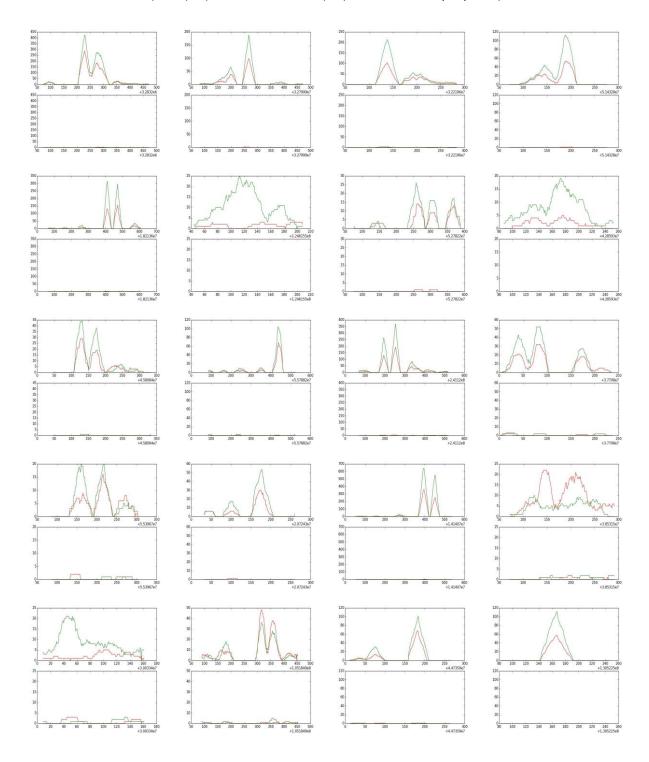
(NRSF) – (PePr vs ZINBA-SA) – (ZINBA-SA-unique peaks)



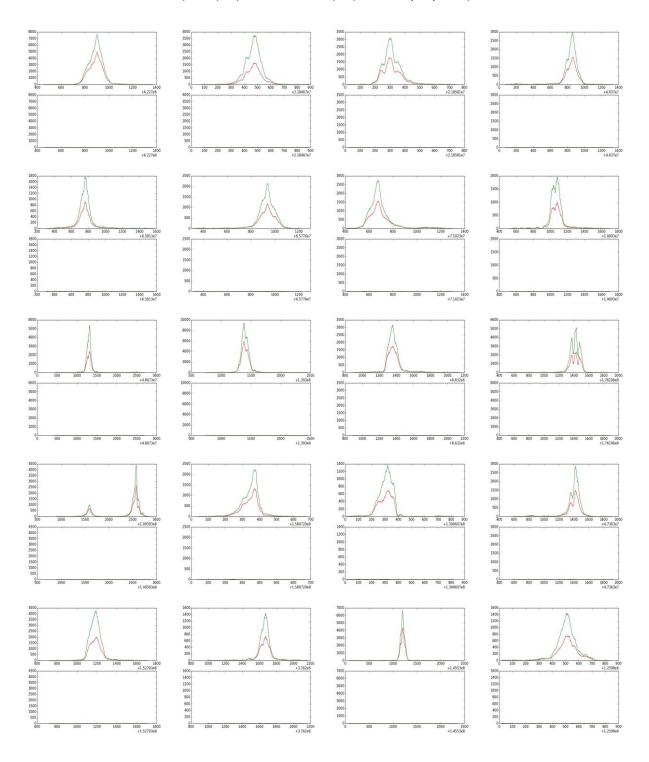
(NRSF) – (PePr vs MACS2-IDR) – (PePr-unique peaks)



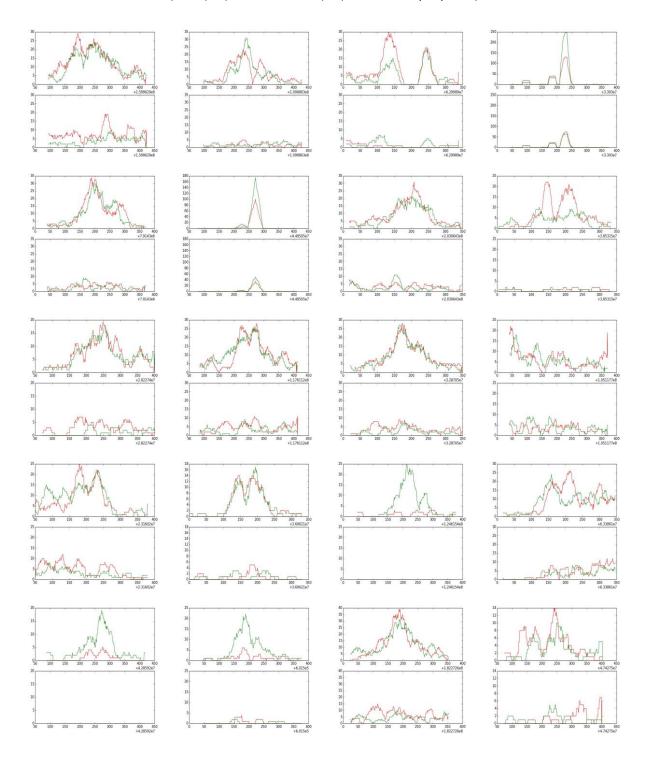
(NRSF) – (PePr vs MACS2-IDR) – (MACS2-IDR-unique peaks)



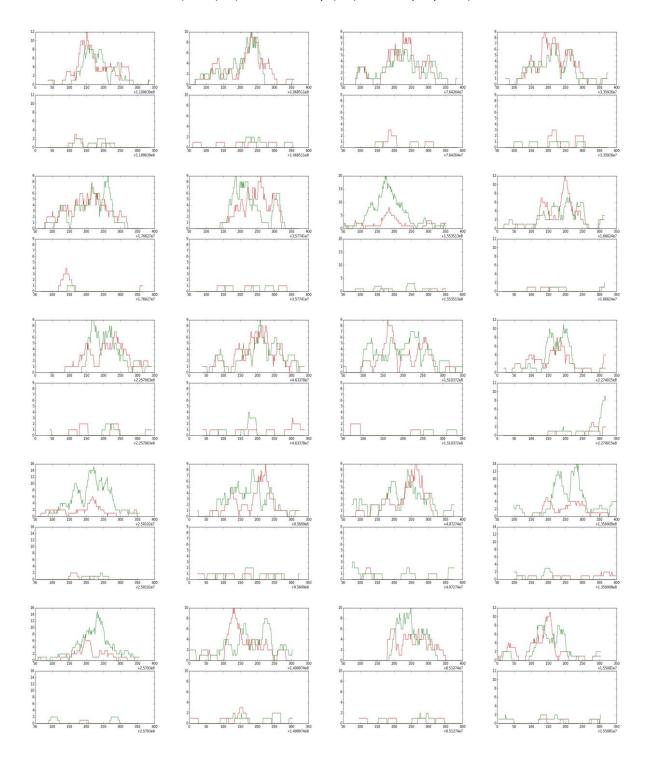
(NRSF) – (PePr vs SPP-IDR) – (PePr-unique peaks)



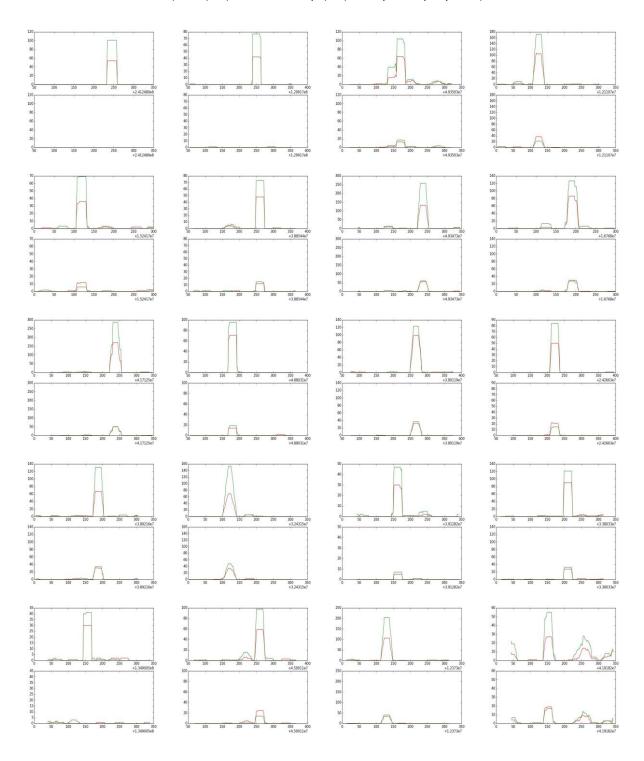
(NRSF) – (PePr vs SPP-IDR) – (SPP-IDR-unique peaks)



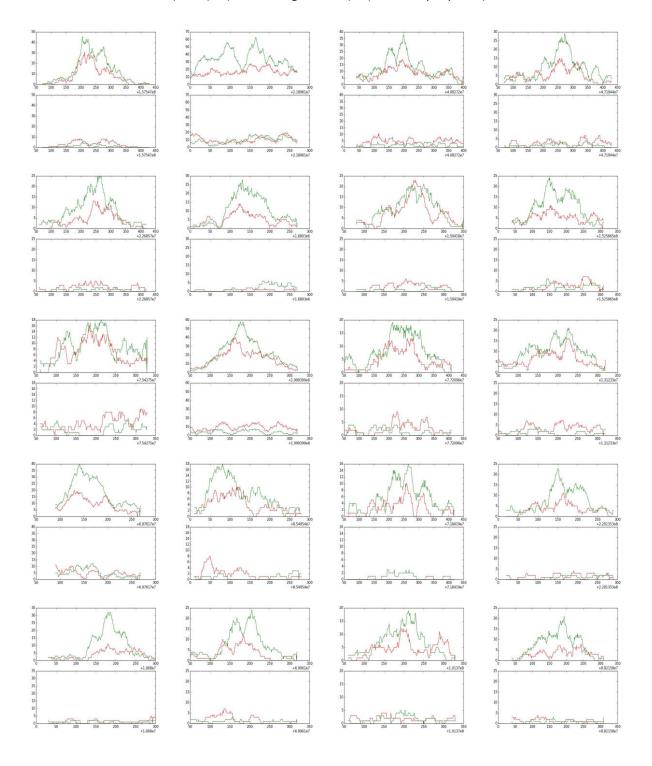
(NRSF) – (PePr vs diffReps) – (PePr-unique peaks)



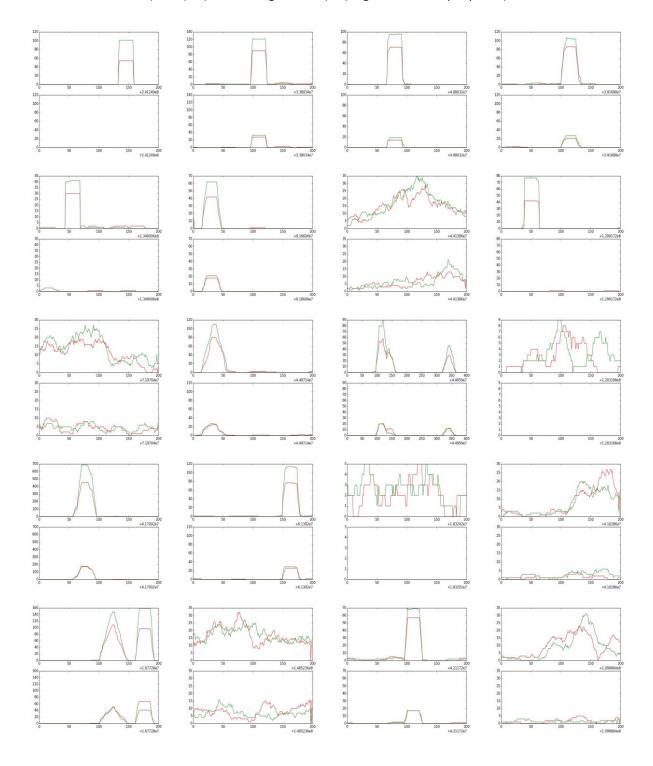
(NRSF) – (PePr vs diffReps) – (diffReps-unique peaks)



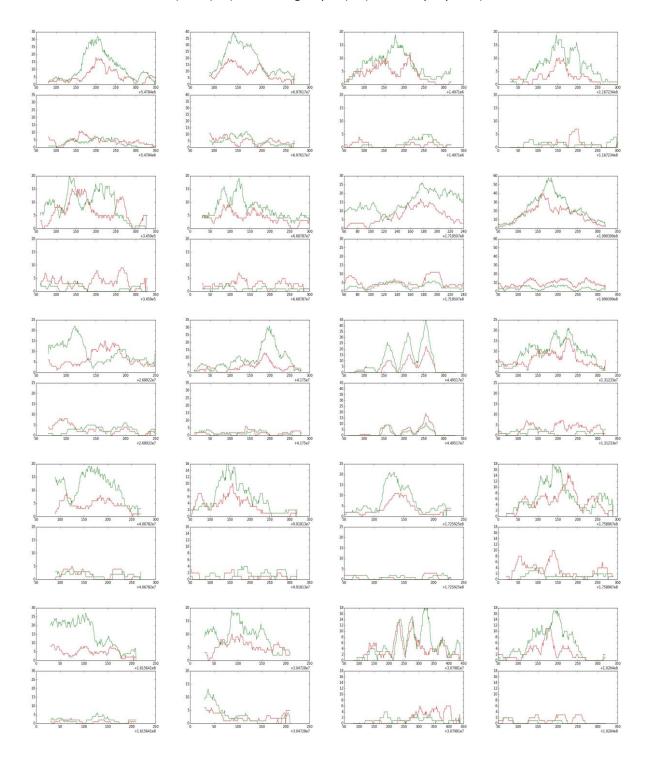
(NRSF) – (PePr vs edgeR-basic) – (PePr-unique peaks)



(NRSF) – (PePr vs edgeR-basic) – (edgeR-basic-unique peaks)



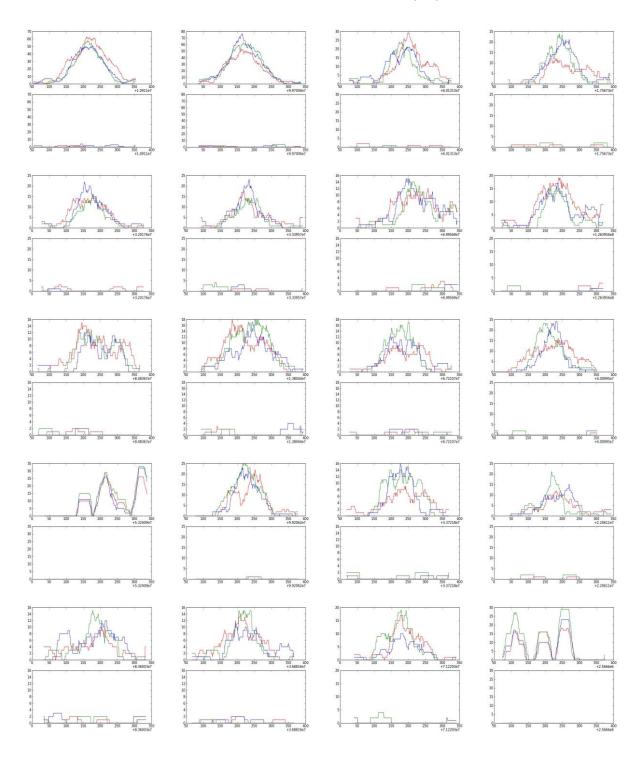
(NRSF) – (PePr vs edgeR-plus) – (PePr-unique peaks)



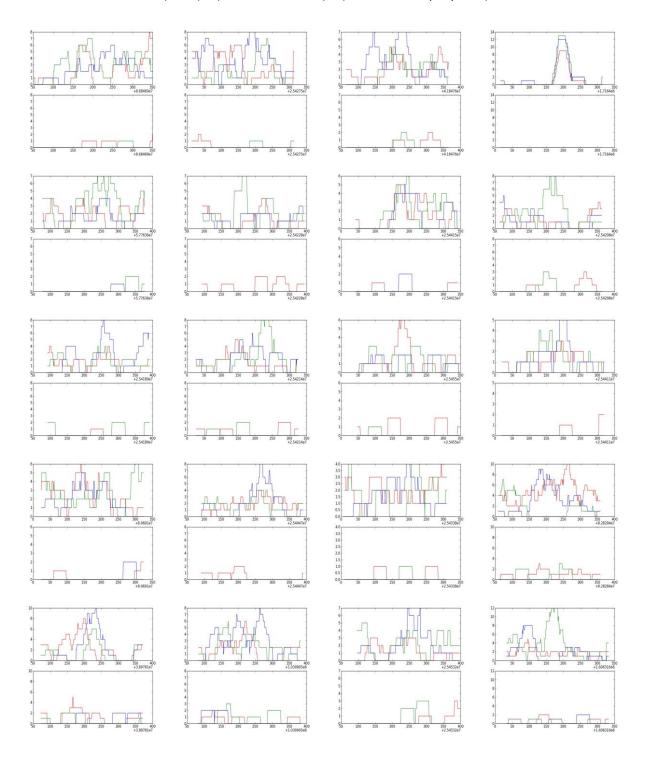
(NRSF) – (PePr vs edgeR-plus) – (edgeR-plus-unique peaks)



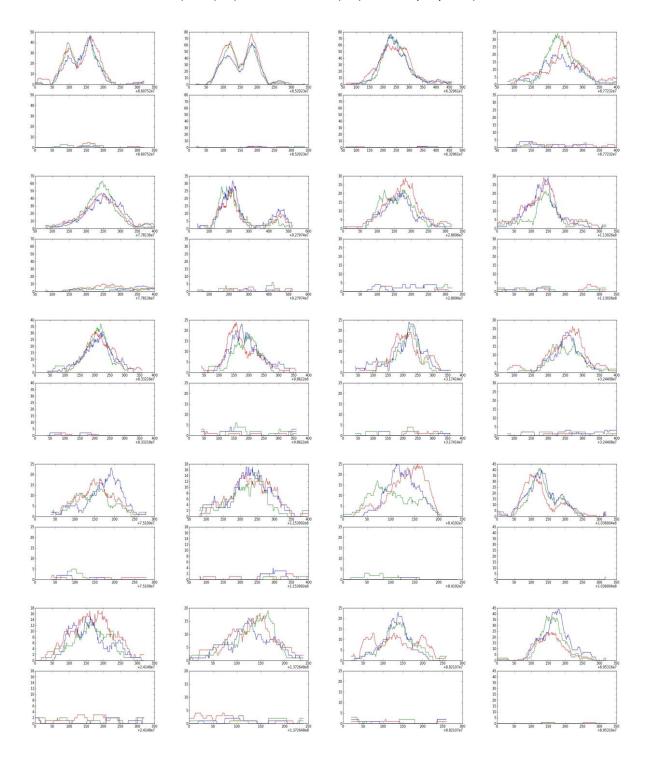
(ATF4) – (PePr vs MACS-CR) – (PePr-unique peaks)



(ATF4) – (PePr vs MACS-CR) – (MACS-CR-unique peaks)



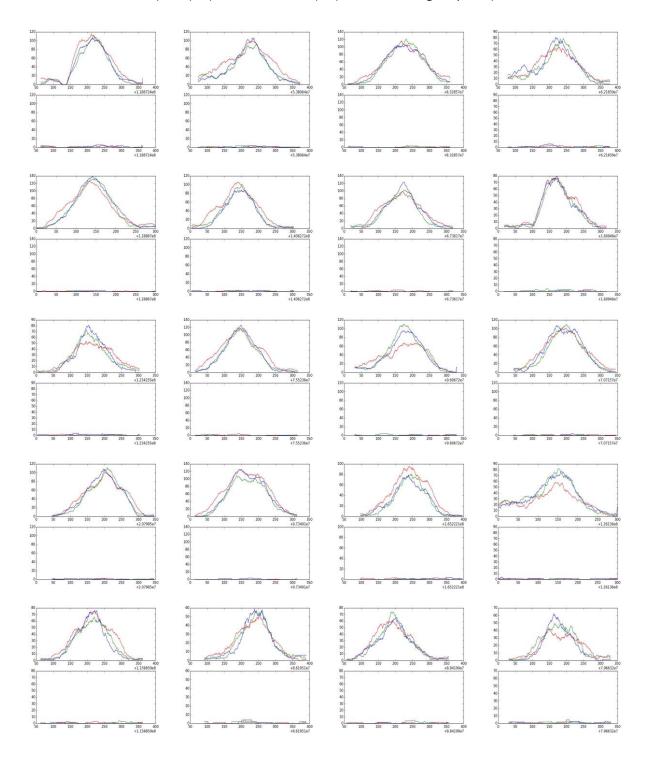
(ATF4) – (PePr vs MACS-SA) – (PePr-unique peaks)



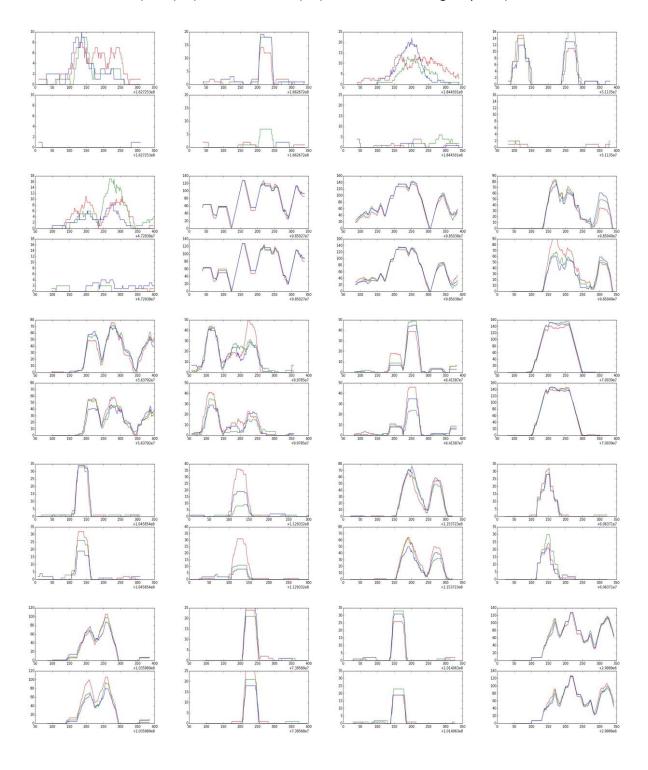
(ATF4) – (PePr vs MACS-SA) – (MACS-SA-unique peaks)



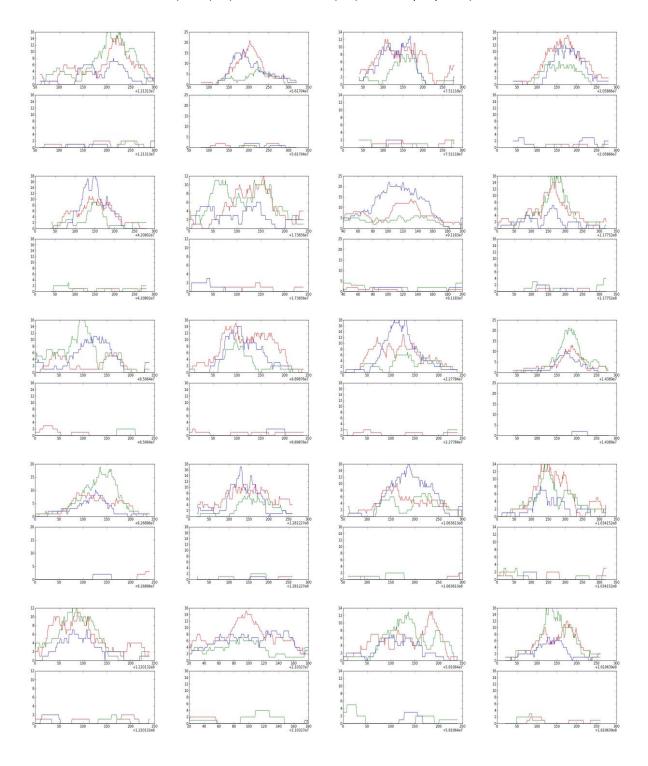
(ATF4) – (PePr vs ZINBA-CR) – (PePr-ranked higher peaks)



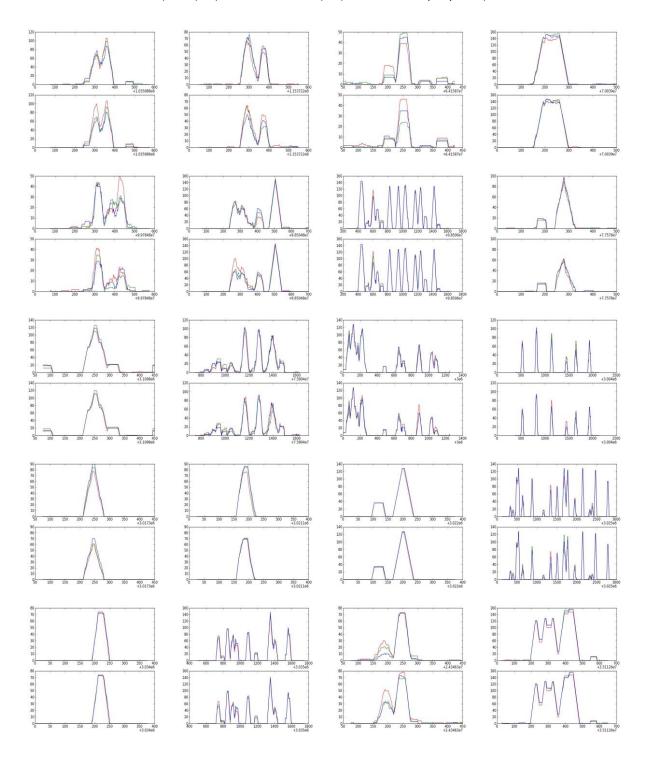
(ATF4) – (PePr vs ZINBA-CR) – (ZINBA-CR-ranked higher peaks)



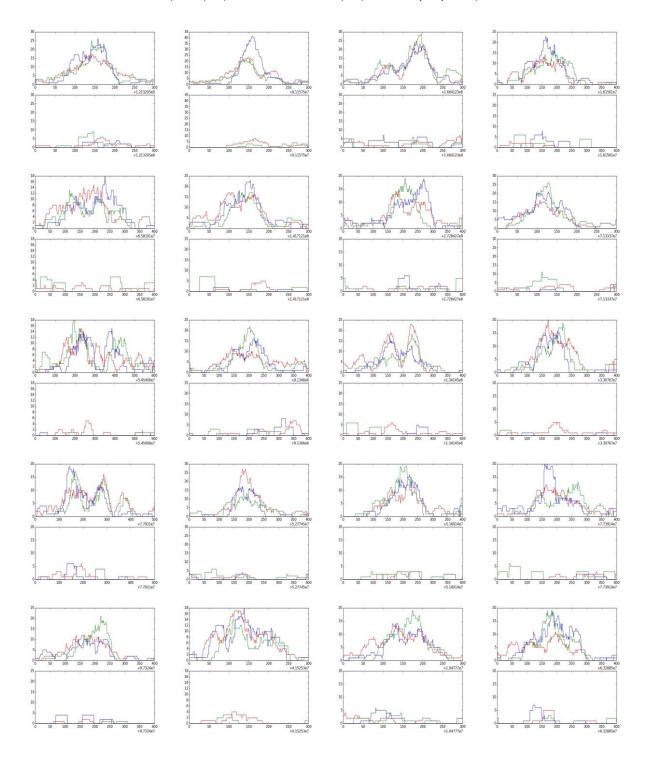
(ATF4) – (PePr vs ZINBA-SA) – (PePr-unique peaks)

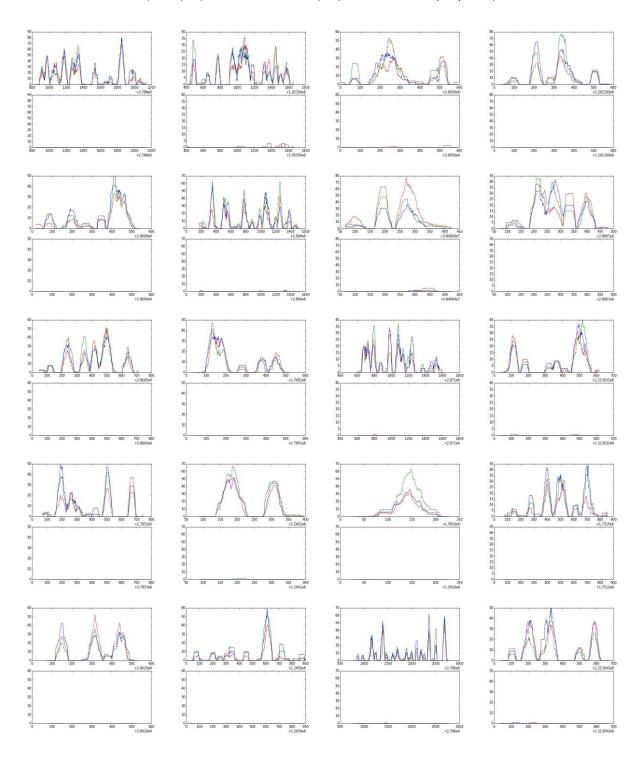


(ATF4) – (PePr vs ZINBA-SA) – (ZINBA-SA-unique peaks)

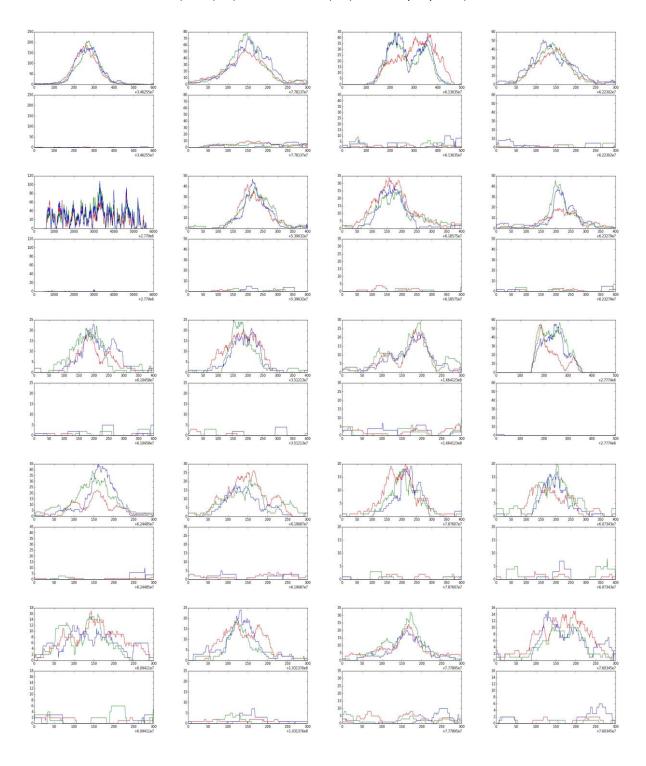


(ATF4) – (PePr vs MACS2-IDR) – (PePr-unique peaks)

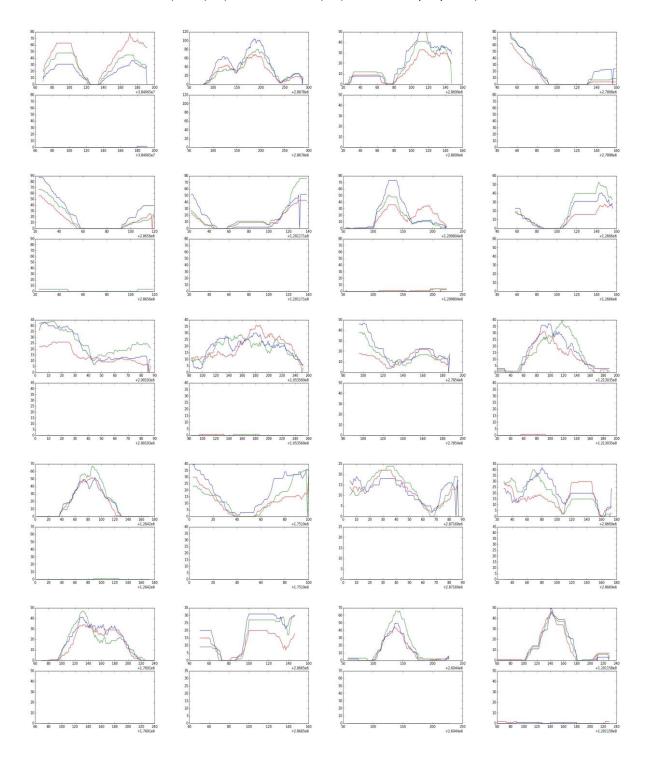




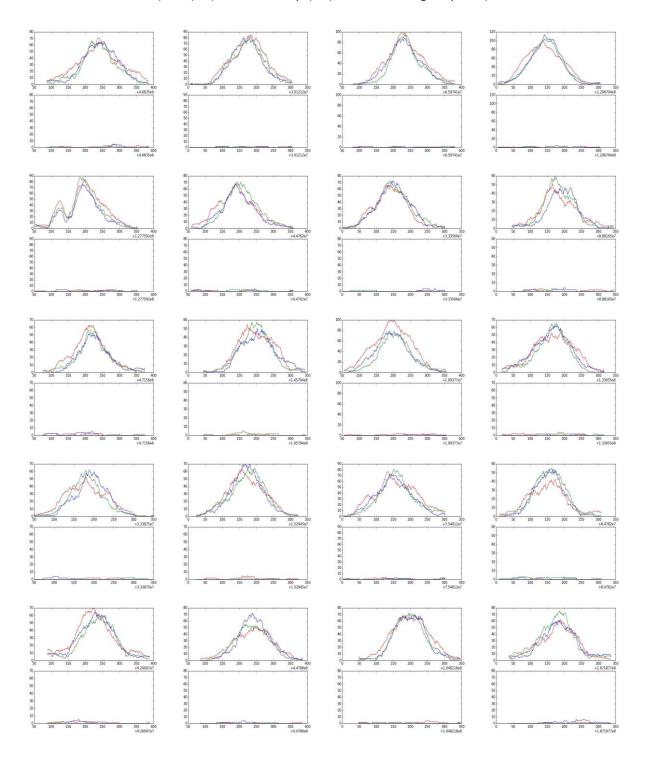
(ATF4) – (PePr vs SPP-IDR) – (PePr-unique peaks)



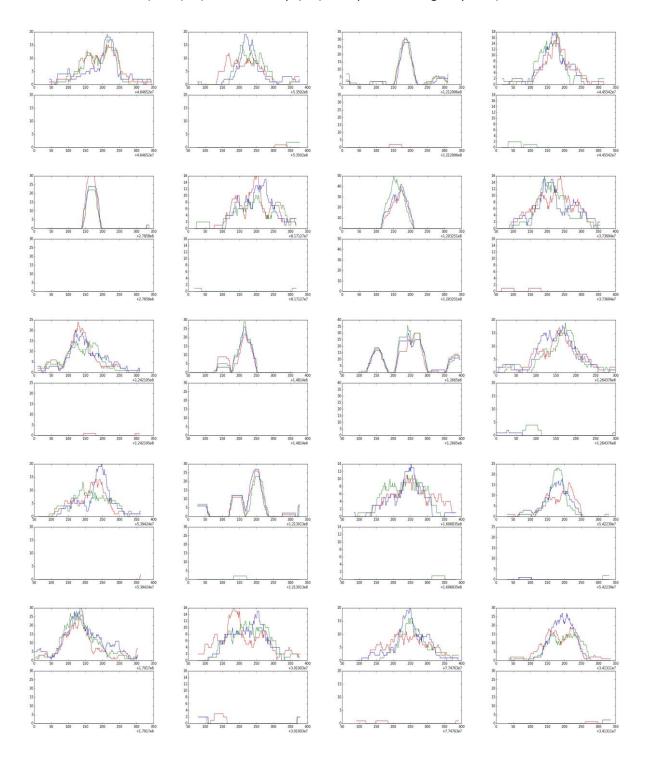
(ATF4) – (PePr vs SPP-IDR) – (SPP-IDR-unique peaks)



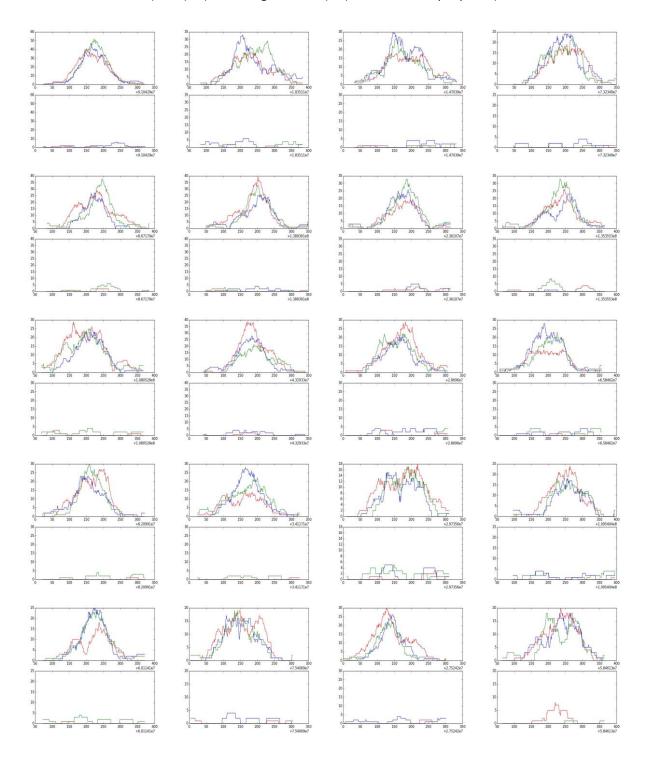
(ATF4) – (PePr vs diffReps) – (PePr-ranked higher peaks)



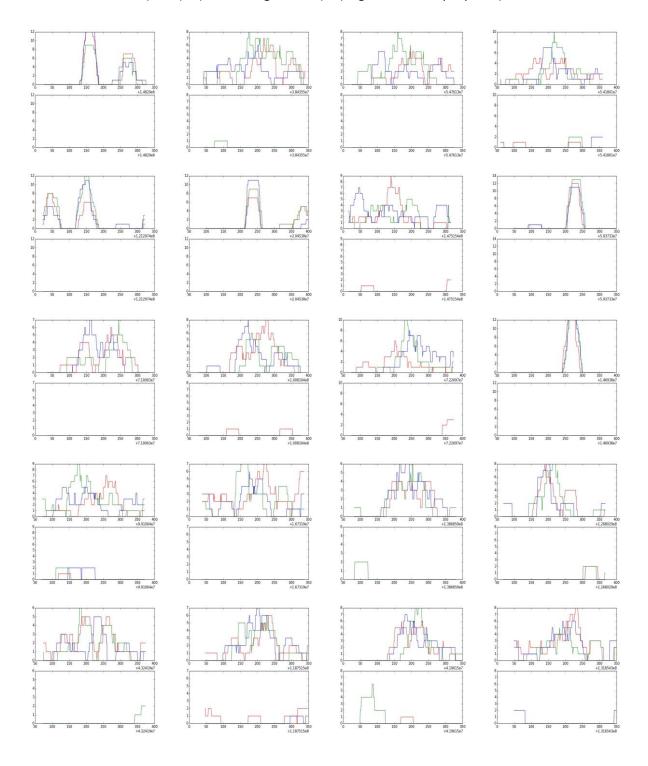
(ATF4) – (PePr vs diffReps) – (diffReps-ranked higher peaks)



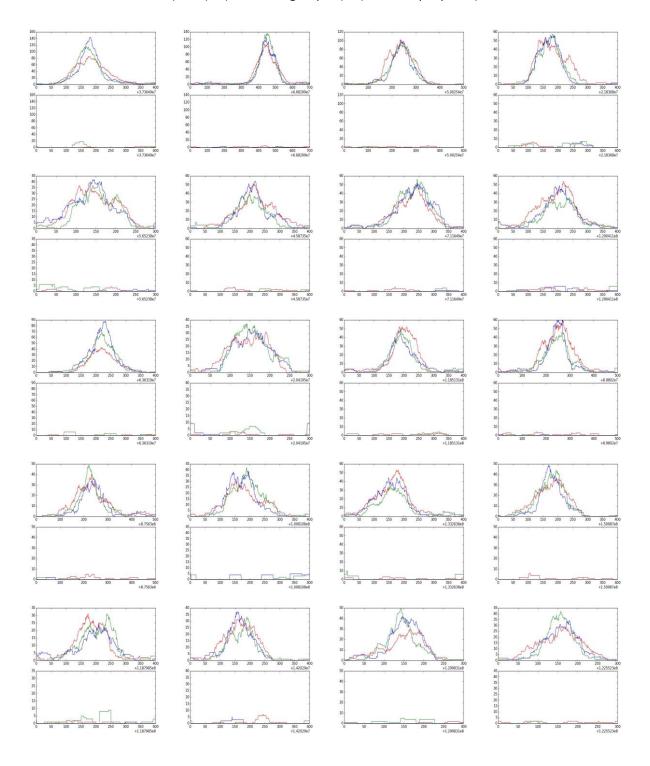
(ATF4) – (PePr vs egdeR-basic) – (PePr-basic-unique peaks)



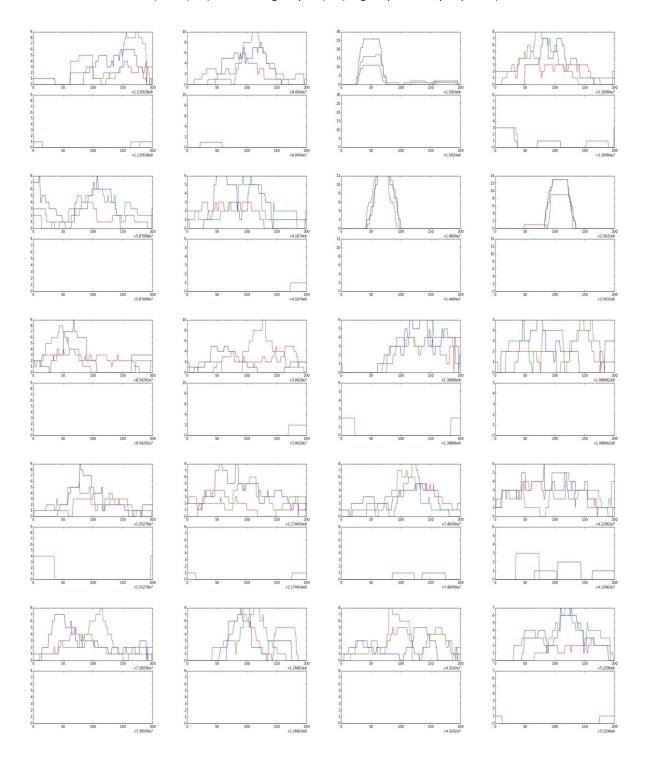
(ATF4) – (PePr vs edgeR-basic) – (edgeR-basic-unique peaks)



(ATF4) – (PePr vs edgeR-plus) – (PePr-unique peaks)



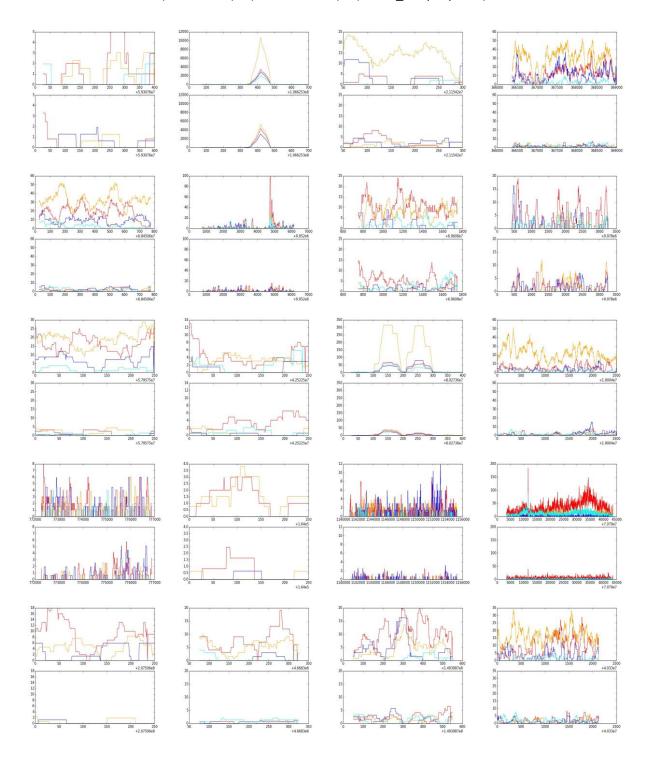
(ATF4) – (PePr vs edgeR-plus) – (edgeR-plus-unique peaks)



(H3K27me3) – (PePr vs ZINBA) – (PePr_unique peaks)



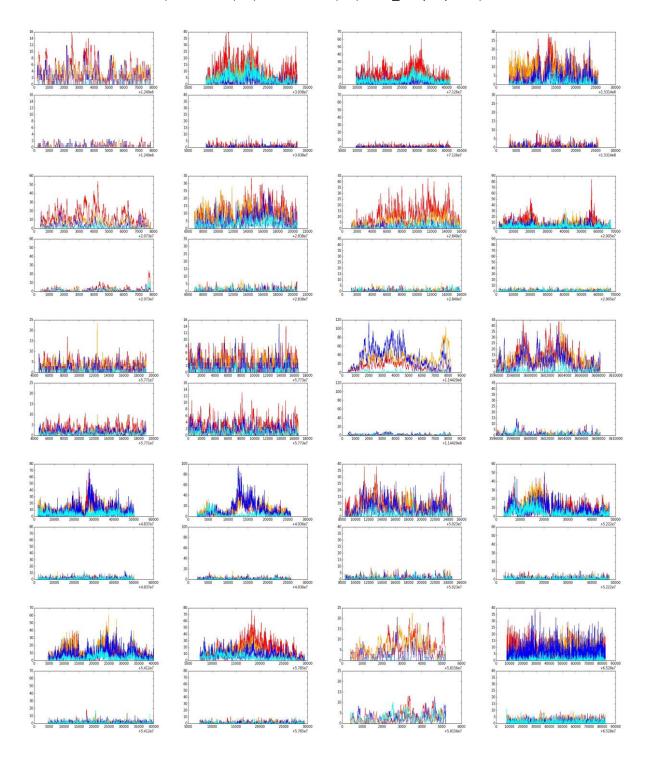
(H3K27me3) – (PePr vs ZINBA) – (ZINBA_unique peaks)



(H3K27me3) – (PePr vs SICER) – (PePr_unique peaks)



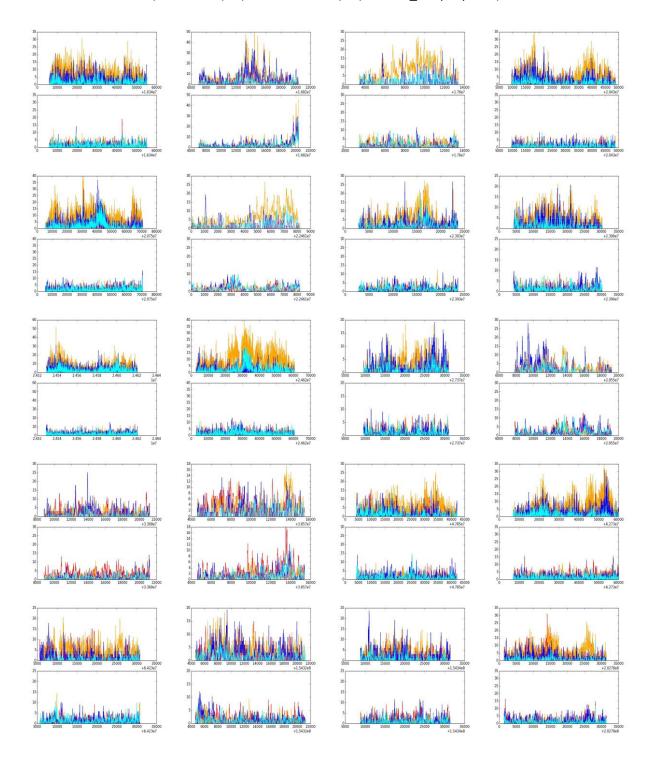
(H3K27me3) – (PePr vs SICER) – (SICER_unique peaks)



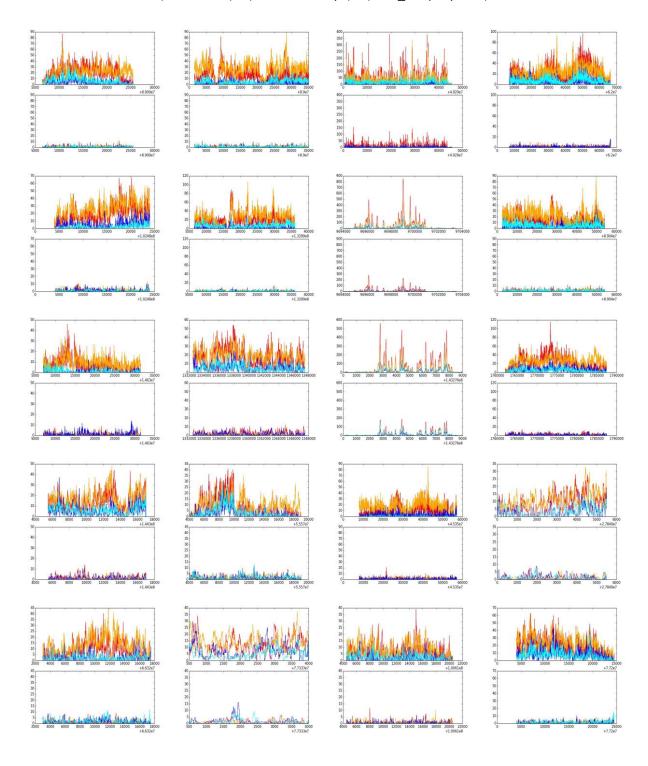
(H3K27me3) – (PePr vs DiffBind) – (PePr_unique peaks)



(H3K27me3) – (PePr vs DiffBind) – (DiffBind_unique peaks)



(H3K27me3) – (PePr vs diffReps) – (PePr_unique peaks)



(H3K27me3) – (PePr vs diffReps) – (diffReps_unique peaks)

